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HH Do, S Melnik, E Rahm - Web, Web-Services, and Database Systems: Node 2002 Web-And ..., 2003 - books.google.com

 ... siveness thus preventing a qualitative **comparison** between systems. ... effort and the hardness of **match** problems. ... very helpful to better **compare** the effectiveness ...

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[A computer vision based technique for 3-D sequence-independent structural comparison of proteins - all 3 versions »](#)

O Bachar, D Fischer, R Nussinov, H Wolfson - Protein Engineering Design and Selection, 2003 - Oxford Univ Press

 ... two typical structure **comparison** problems and **compare** them with ... provides an option to allow self **comparison**, which produces the best **match** besides die ...

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[Large-Scale Comparison of Intron Positions among Animal, Plant, and Fungal Genes - all 8 versions »](#)

A Fedorov, AF Merican, W Gilbert - Proceedings of the National Academy of Sciences of the ..., 2002 - National Acad Sciences

 ... alignment to **compare** these 30 genes to fungal-intron ... introns from the best fungal-animal **match** onto the ... such an animal-plant-fungal **comparison**, and **Table** ...

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 A Krogh^f, I Brown - J. Mol. Biol., 1994 - cab.zju.edu.cn

 ... of an HMM are estimated from a training **set** of unaligned sequences. ... To demonstrate the method, we examine three protein families. Each family consists of a **set** ...

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MJ Wise - Proceedings of the twenty-seventh SIGCSE technical symposium ..., 1996 - portal.acm.org

 ... the problem exists of finding the initial **set** of unique ... to reduce the otherwise $O(n^2)$ cost of this **comparison**. ... of maximal-matches length L do **remove match**(p, t ...

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[Learning to **match** and cluster large high-dimensional data sets for data integration - all 10 versions »](#)

WW Cohen, J Richman - Proceedings of the eighth ACM SIGKDD international ..., 2002 - portal.acm.org

 ... and • in Step 3d, let $T_{t,i} = 0$ (ie, only **remove** a from the **set** of PossibleCenters ...

 Two of the datasets require clustering, and two require **match**- ing ...

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[S-Match: an Algorithm and an Implementation of Semantic Matching - all 18 versions »](#)

F Giunchiglia, P Shvaiko, M Yatskevich - The Semantic Web: Research and Applications: First European ..., 2004 - books.google.com

 ... step 3) and, finally (step 4), we **compare** C_2 in ... Evaluation We have done some preliminary **comparison** between S ... is an estimate of the post **match** efforts needed ...

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LP Cox, CD Murray, BD Noble - ACM SIGOPS Operating Systems Review, 2002 - portal.acm.org

 ... meta-data is not chunked and is mu- **table**. ... with commonly-appearing chunkids, hoping for a false **match**. ... We call these the add **set**, delete **set**, and meta-data ...

[Cited by 150](#) - [Related Articles](#) - [Web Search](#)
[Fast algorithms for large-scale genome alignment and comparison - all 33 versions »](#)

AL Deicher, A Phillippy, J Carlton, SL Salzberg - Nucleic Acids Research, 2002 - Oxford Univ Press

 ... $4k + 1$ bytes for a minimum **match** size of ... algorithmic extensions that permit alignment and **comparison** of protein ... It has been used to **compare** sequences as long ...

[Cited by 228](#) - [Related Articles](#) - [Web Search](#)

[PDF] [Automatic grammar induction and parsing free text: A transformation-based approach](#) - [all 20 versions ...](#)

E Brill - [Proceedings of the 31st Annual Meeting of the Association ...](#), 1993 - [acl.ldc.upenn.edu](#)

... presented above allow us to **compare** the transformation ... carried out are integer addition and integer **comparison**. ... hope that this grammar will **match** the grammar ...

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